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Figure 1

-150 -TGGGCTCAGCCACGCCCCAGGGTCCCCCAGTGGGACTAGTTCTTCATTCTGCGAGCTGCACACATCTGTCACTGAGGGAATGTCAAGTC
-60 -TCTCACTCTCCTCTCCTCACTATCTTTCCGAGAAAGCGGGTCTCTCTGCTTGTGCGAGTATGGACGACCCGGACTGCGATTCCACCTGG
MetAspAspProAspCysAspSerThrTrp 10

31 -CAGCAGGAGAGCGAGGAGCATGCCGAGGATGCCAGGCGGATGATACGACCGATGAGGACACGGGGCAGCATGACGGCGACGGCGAGGAG
GluGluGluSerGluGluAspGlyGluAspGlyGlnAlaAspAspThrThrAspGluAspThrGlyAspAspAspGlyAspAlaGluGlu 40

121 -GCACGGCCAAGCCTGTTCCAGTCCAGGATGACAGCGGTACCGAACTGCGGTCTATGCAGGACATGCAAGATACCGGCACAACCTACCCG
AlaArgProSerLeuPheGlnSerArgMetThrGlyTyrArgAsnTrpArgAlaMetGlnAspMetGlnArgTyrArgHisAsnTyrPro 70

211 -GATTTGACAGATCAAGACTGCAATGGTGACATGTGCAACCTGAGCTTCTACAAAAATGAGATCTGCTTCCAGCCAAATGGGGCTCTCATC
AspLeuThrAspGlnAspCysAsnGlyAspMetCysAsnLeuSerPheTyrLysAsnGluIleCysPheGlnProAsnGlyAlaLeuIle 100

301 -CAGGACATTCTTCAGAACTGCAAGACAACCTATGACCTCCTGCAAGAGAATCACTCCTACATCCAGTGGCTCTTCTCTCGCGGAACCA
GluAspIleLeuGlnAsnTrpLysAspAsnTyrAspLeuLeuGluGluAsnHisSerTyrIleGlnTrpLeuPheProLeuArgGluPro 130

391 -GGAGTGAAGTGGCAGCCCAAGCCCTCACCTGCAAGGAGGTTGAGGCATTTAAAGCTCCAAGGAAGTCAGAGAGCGTCTTGTCCGGGCC
GlyValAsnTrpHisAlaLysProLeuThrLeuLysGluValGluAlaPheLysSerSerLysGluValArgGluArgLeuValArgAla 160

481 -TATGAGCTCATGCTGGGCTTCTATGGGTTCACCTTGAGGACCGGGGCACGGGTGCTGTATGCCGTGCACAGAACTTCCAGCCCGGCTTC
TyrGluLeuMetLeuGlyPheTyrGlyPheHisLeuGluAspArgGlyThrGlyAlaValCysArgAlaGlnAsnPheGlnProArgPhe 190

571 -CACAACTCTGAACAGCCACAGCCACAACCTGCGTATTACAGCATCTCTCAAGTCACTGGGTGAGCTGGGCTTAGAACACTACCAGGCA
HisAsnLeuAsnSerHisSerHisAsnAsnLeuArgIleThrArgIleLeuLysSerLeuGlyGluLeuGlyLeuGluHisTyrGlnAla 220

661 -CCCCGGTCCGCTTCTTCTCGAGGAGACCCCTTGTACAGCACAACCTGCCACGGTGGCCAGAGTGGCCCTGGACTACTTCTGTTCGGT
ProLeuValArgPhePheLeuGluGluThrLeuValGlnHisLysLeuProSerValArgGlnSerAlaLeuAspTyrPheLeuPheAla 250

751 -GTGCGCTGCGGGCACCAGCGCGGGAGCTTGTGTACTTTGCTTGGGAGCACTTCAAGCTCGCCGAGAGTTGTCTGGGGGCCCCGTGAC
ValArgCysArgHisGlnArgArgGluLeuValTyrPheAlaTrpGluHisPheLysProArgArgGluPheValTrpGlyProArgAsp 280

841 -AAGCTGCGGAGATTCAAGCCCCAGACCATACCCACGCCACTGACGGGACCCAGGGCAGGATAAAGATGAGGGCTCCAGGGACCCCTCC
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931 -AAGAGCCTGCGACCCAGCGTCCGACCTGTGGATCTGCAAGGGACCTGAGTGGGACAGTGGAACAGCTGAGGATCCCTCACTGCTGAAC
GlnGluAlaGlyThrGlnGlyArgThrCysGlySerGlyArgAspLeuSerGlyAspSerGlyThrAlaGluAspProSerLeuLeuAsn 340

1021 -ACAAAGCCCTCAGATGGGGGAACCTTGGATGGGAACCAGAGGATGAAGCTAAGTCCCTGAGTCCCAAGGAGAGCAAGAAAAGGAAGTTG
ThrLysProSerAspGlyGlyThrLeuAspGlyAsnGlnArgAspGluAlaLysSerLeuSerProLysGluSerLysLysArgLysLeu 370

1111 -GAGGGGAACAGGCAGGAGCAGGTCCACGGGAGGCGAGATCCCCAGGGTGTCTCTGAGGTAGAGAAAATTGCCCTTAACCTTGAGGAGTGT
GluGlyAsnArgGlnGluGlnValProGlyGluAlaAspProGlnGlyValSerGluValGluLysIleAlaLeuAsnLeuGluGluCys 400

1201 -GCCCTTAGCCCTATCAGCCAGGAGCCAGGGAGGCTGAACCGCCCTGTCTGTGGCCAGGCTGGCTAATGAGGTAAGAAAGCGGACGAG
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1291 -GTGGAGGAAGGGGCTGAGGGTATGGAGTAGTCAGTAACACTCAAATGCAGGCCAGTGCCTGCCTCCTACCCCTTCAGAGTGTCTCTGAG
ValGluGluGlyAlaGluGlyAspGlyValValSerAsnThrGlnMetGlnAlaSerAlaLeuProProThrProSerGluCysProGlu 460

1381 -GCCCCAAAGGATGGGAATGGGCCAGAGGACTCAAACAGCCAGGTTGGGGCAGAGGATTCCAAAGCCAGGTGGGGCCGGAGGATCCAAAC
AlaGlnLysAspGlyAsnGlyProGluAspSerAsnSerGlnValGlyAlaGluAspSerLysSerGlnValGlyProGluAspProAsn 490

1471 -AGCCAGGTGGGGCTGGAGGACCCAAACAGCCAGGTGGGGCAGAGGACCCAAACAGCCAGGTGGGGCAGAGGACCCAAACAGCCAGGTG
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1561 -GGGCCAGAGGACCCAAACAGCCAGGTGGGGCAGAGGACCCAAACAGCCAGGTGGTGGGGCAGAGCAAGCTGCCTCTAAGAGCCCTGTG
GlyProGluAspProAsnSerGlnValGlyProGluAspProAsnSerGlnValGlyProGluGlnAlaAlaSerLysSerProVal 550

1651 -GAGGACCTGACTCTGACACTATGGGAACCTCAGTGGATGAGTCAGAGGAGTTGGCAAGGATTGAGGCTCTGTGAACCCCCAAAGCCT
GluAspProAspSerAspThrMetGlyThrSerValAspGluSerGluGluLeuAlaArgIleGluAlaSerAlaGluProProLysPro 580

1741 -TAGAGGTGCATCTCAGTCTACTCAGCCCACTGCAGGGGGTTTCTGAGTCCAGAGCTCTGCCGTAGGCTCTTCTTGGTGGCCACAGTGC
1831 -TGGCCTCTCCCTAGTGGTCACTGAGGTGGCCACCAGAGGACTGAGGCCCTGCCCTCAGGGAAGGCCAAGGCCCTTCAGAACCCCTCTTAC
1921 -CTCACTGTGTCTCTCTCACTGCCCTCTGAGCCCTGCCCTTGTGATCAGACCTTAAGGGTCTAGAGGGAGGGGCTCTTCATTAGTCTGCT
2011 -GCCAAGTGAGGCGCTTTTCTGAATAAACTCTTTAGACTTTGTCAA

Figure 2

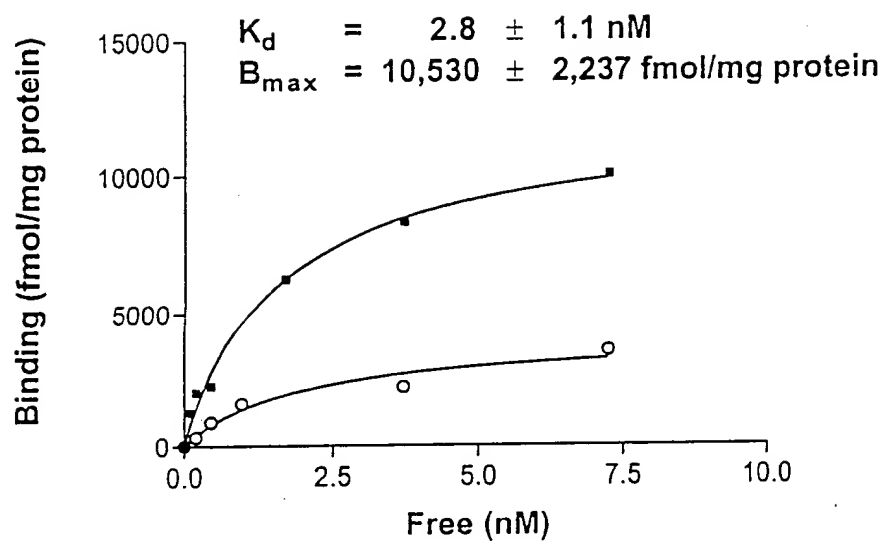


Figure 3

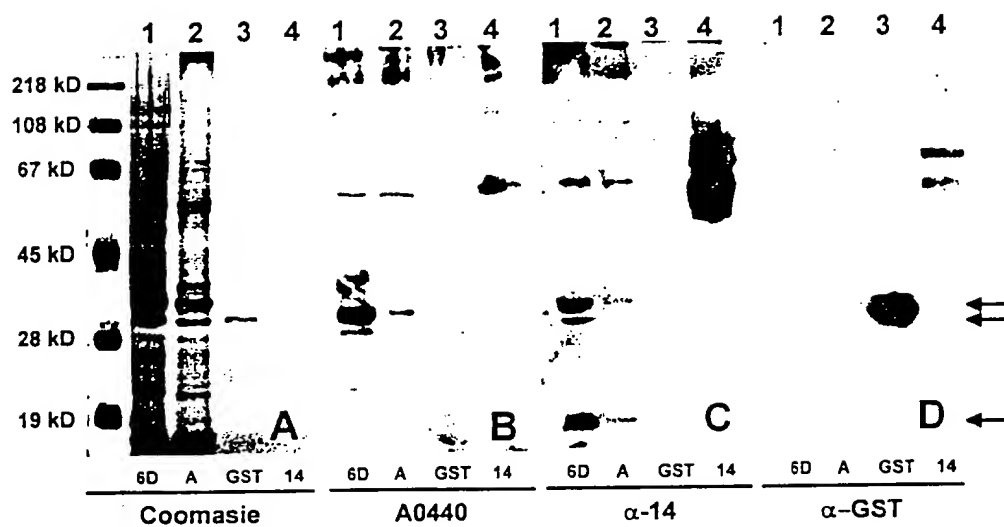


Figure 4

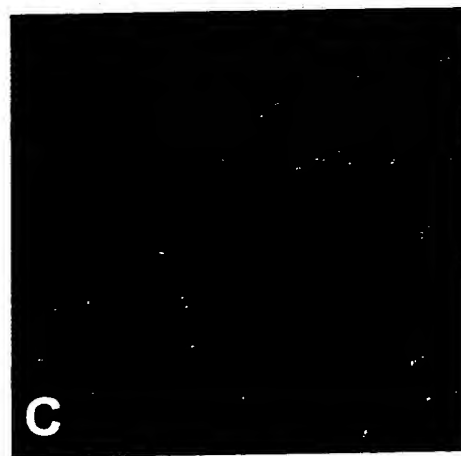
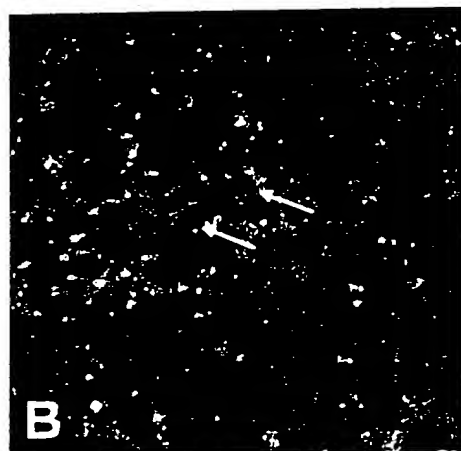
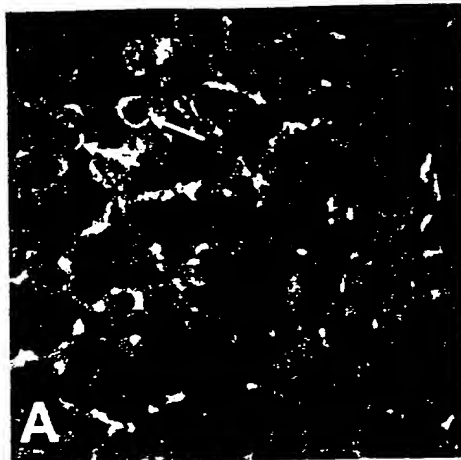


Figure 5

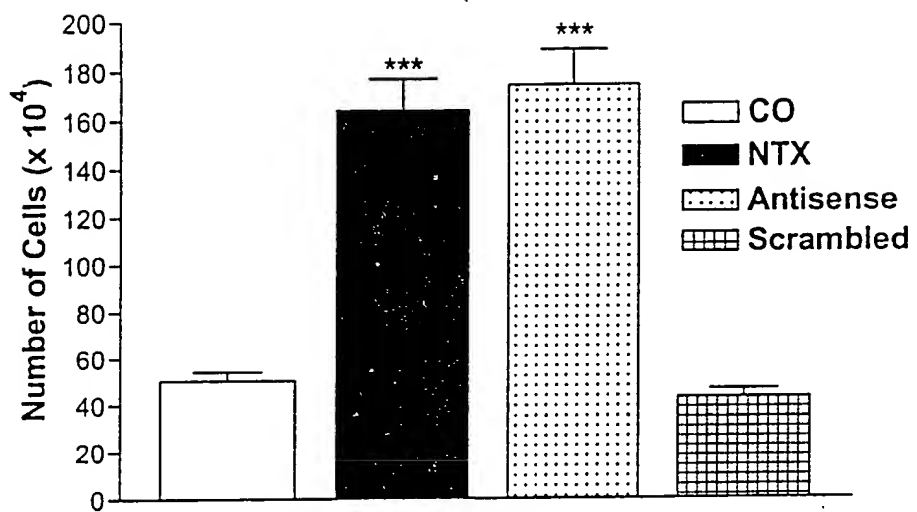


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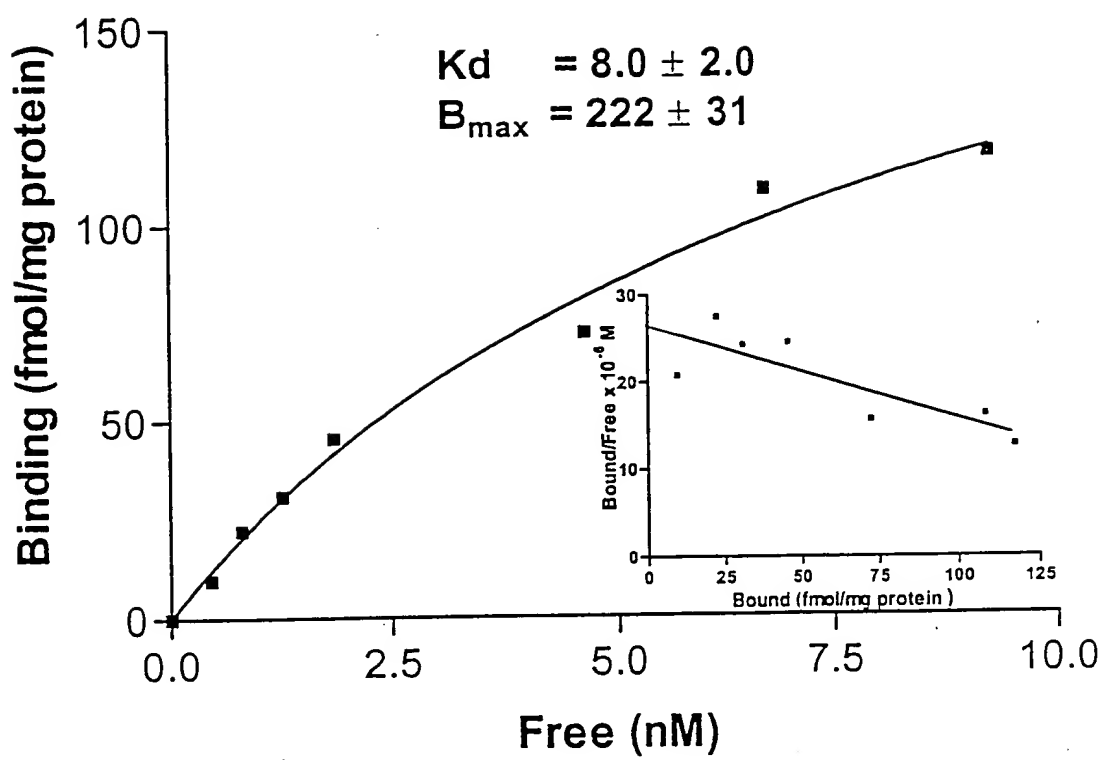


Figure 7

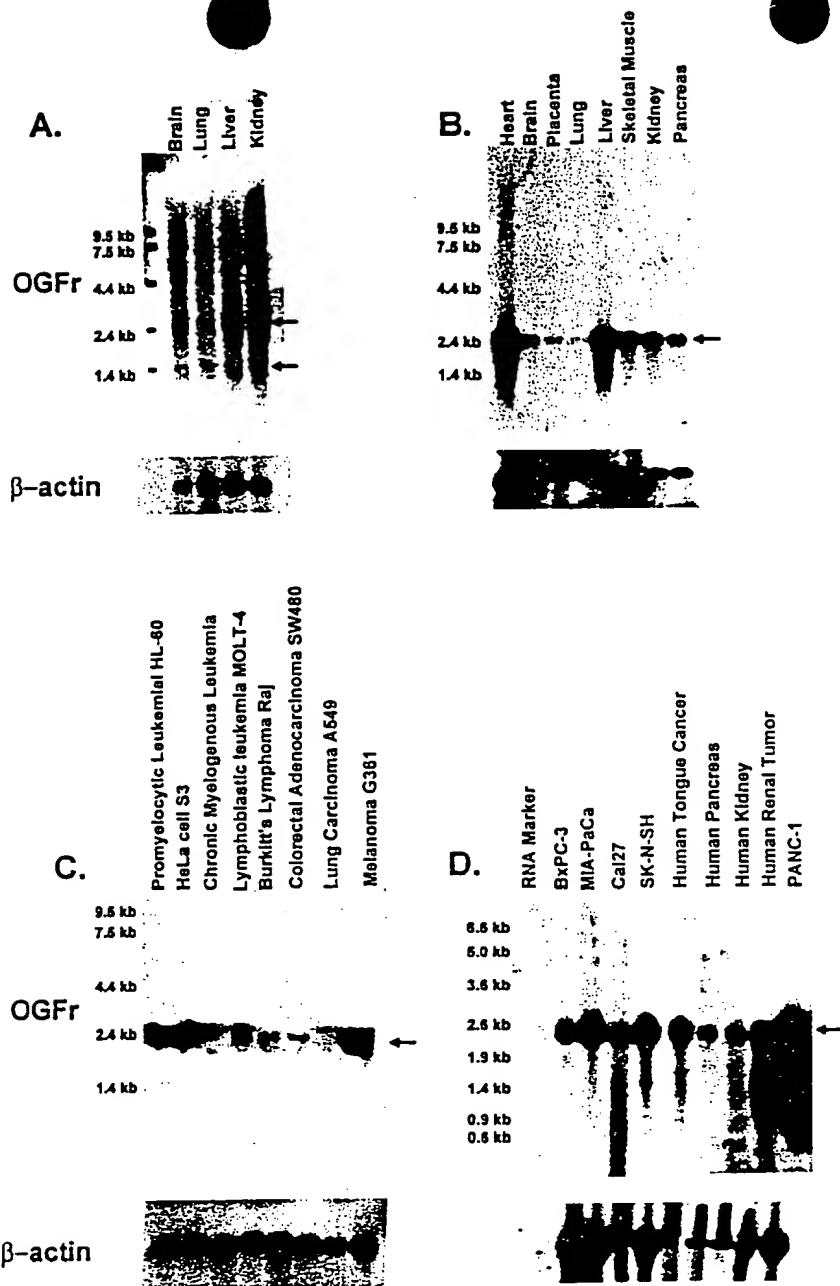


Figure 9

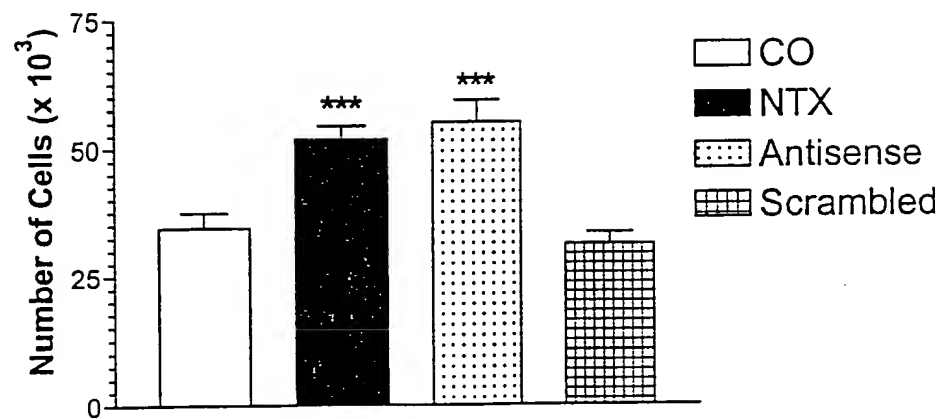


Figure 10

Human and Rat OGFr

79% Identical/ 87% Similar	39.5% Identical/ 56% Similar	23% Identical/ 47% Similar	20% Identical/ 43% Similar	
1	297	464	629	697
Amino Acid Number				

Figure 11

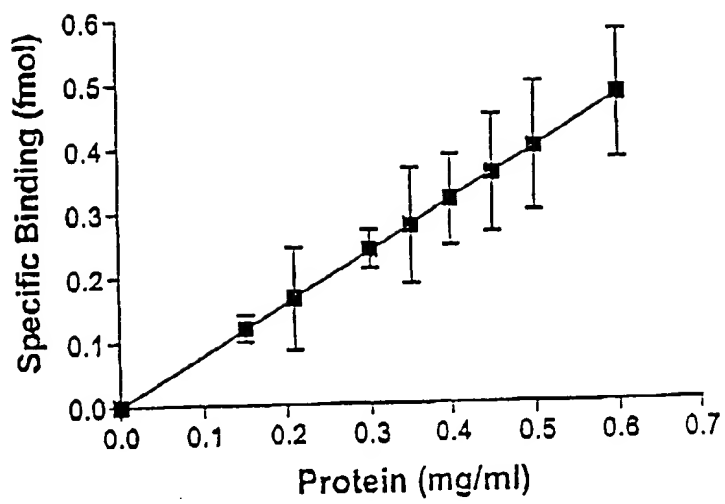


Figure 12

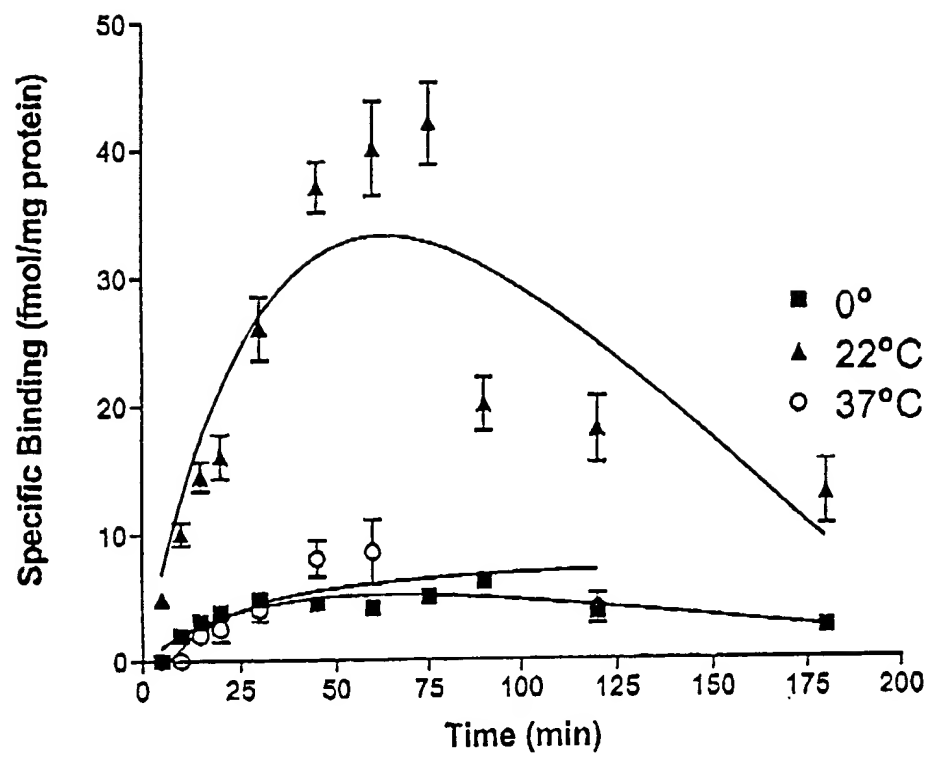


Figure 13

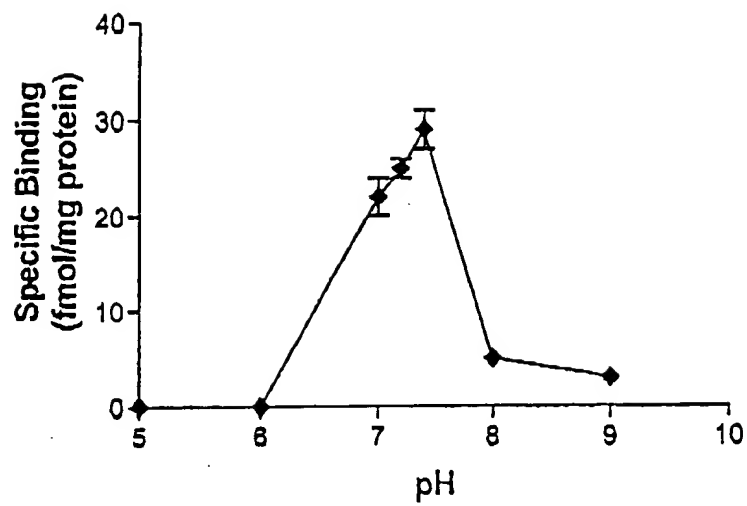


Figure 14

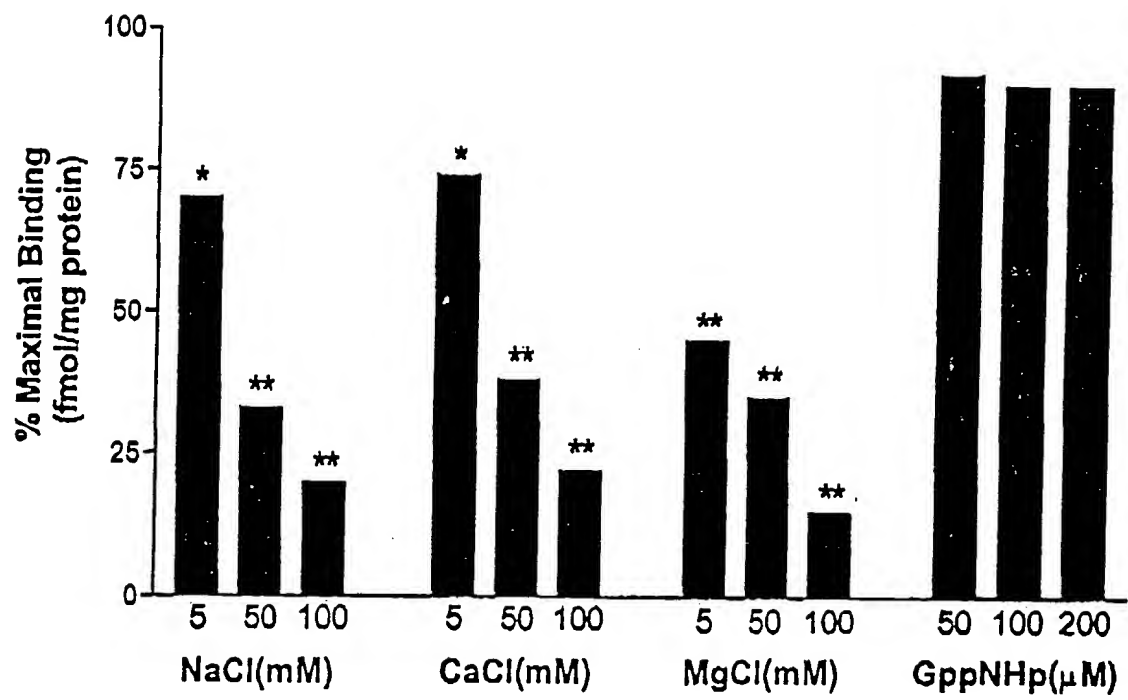


Figure 15

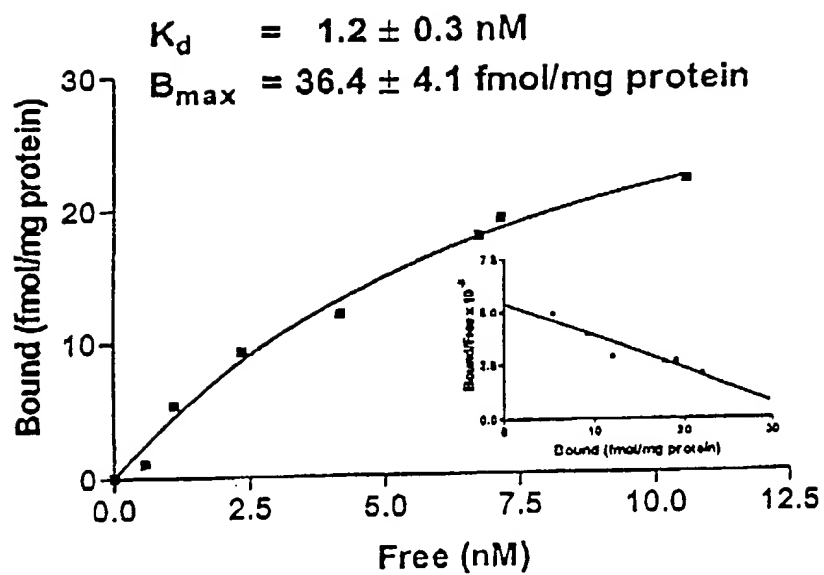


Figure 16